



PCT

RAW SEQUENCE LISTING

DATE: 05/16/2005

PATENT APPLICATION: US/10/534,135

TIME: 12:10:05

Input Set : A:\PTO.SR.txt

Output Set: N:\CRF4\05162005\J534135.raw

3 <110> APPLICANT: Johnson & Johnson Research Pty Ltd
W--> 4 <120> TITLE OF INVENTION: A means of producing and utilising a population of disease
W--> 5 specific cytotoxic T lymphocytes
W--> 6 <130> FILE REFERENCE: 02 1355 7401
C--> 7 <140> CURRENT APPLICATION NUMBER: US/10/534,135
C--> 7 <141> CURRENT FILING DATE: 2005-05-06
W--> 7 <160> NUMBER OF SEQ ID: 9
8 <170> SOFTWARE: PatentIn version 3.3

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed
(Pg. 1)

86 <210> SEQ ID NO: 9
87 <211> LENGTH: 76
88 <212> TYPE: DNA
89 <213> ORGANISM: Artificial
W--> 90 <220> FEATURE:
91 <223> OTHER INFORMATION: 3L1Dz5 primer
W--> 92 <400> SEQUENCE: 9
93 caccaaaaga gaactgcaat tcgttgtagc tagcctttca ggaccacag gagcggcaag 60
95 caattcgttc tgtatc 76
E--> 102 73.
E--> 104 141719585
E--> 105 141719585

Deleted

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/534,135

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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,6,7,8,9

VERIFICATION SUMMARY

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L:4 M:283 W: Missing Blank Line separator, <120> field identifier
L:6 M:283 W: Missing Blank Line separator, <130> field identifier
L:7 M:270 C: Current Application Number differs, Replaced Current Application No
L:7 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:7 M:283 W: Missing Blank Line separator, <160> field identifier
L:14 M:283 W: Missing Blank Line separator, <400> field identifier
L:21 M:283 W: Missing Blank Line separator, <400> field identifier
L:28 M:283 W: Missing Blank Line separator, <220> field identifier
L:30 M:283 W: Missing Blank Line separator, <400> field identifier
L:37 M:283 W: Missing Blank Line separator, <400> field identifier
L:46 M:283 W: Missing Blank Line separator, <400> field identifier
L:54 M:283 W: Missing Blank Line separator, <220> field identifier
L:56 M:283 W: Missing Blank Line separator, <400> field identifier
L:64 M:283 W: Missing Blank Line separator, <220> field identifier
L:66 M:283 W: Missing Blank Line separator, <220> field identifier
L:70 M:283 W: Missing Blank Line separator, <220> field identifier
L:74 M:283 W: Missing Blank Line separator, <400> field identifier
L:81 M:283 W: Missing Blank Line separator, <220> field identifier
L:83 M:283 W: Missing Blank Line separator, <400> field identifier
L:90 M:283 W: Missing Blank Line separator, <220> field identifier
L:92 M:283 W: Missing Blank Line separator, <400> field identifier
L:102 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:77 SEQ:9
L:102 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
M:254 Repeated in SeqNo=9
L:105 M:252 E: No. of Seq. differs, <211> LENGTH:Input:76 Found:77 SEQ:9